

OIPF

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/717,641  
 DATE: 12/06/2000  
 TIME: 15:13:14

Input Set : A:\P0645P4D2C3.txt  
 Output Set: N:\CRF3\12062000\I717641.raw

## SEQUENCE LISTING

**ENTERED**

## W--&gt; 3 SEQUENCE LISTING

## 5 (1) GENERAL INFORMATION:

7 (i) APPLICANT: Garrard, Lisa J.  
 8 Henner, Dennis J.  
 9 Bass, Steven  
 10 Greene, Ronald  
 11 Lowman, Henry B.  
 12 Wells, James A.  
 13 Matthews, David J.

15 (ii) TITLE OF INVENTION: ENRICHMENT METHOD FOR VARIANT PROTEINS WITH  
 16 ALTERED BINDING PROPERTIES

18 (iii) NUMBER OF SEQUENCES: 43

20 (iv) CORRESPONDENCE ADDRESS:

21 (A) ADDRESSEE: Genentech, Inc.  
 22 (B) STREET: 1 DNA Way  
 23 (C) CITY: South San Francisco  
 24 (D) STATE: California  
 25 (E) COUNTRY: USA  
 26 (F) ZIP: 94080

28 (v) COMPUTER READABLE FORM:

29 (A) MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk  
 30 (B) COMPUTER: IBM PC compatible  
 31 (C) OPERATING SYSTEM: PC-DOS/MS-DOS  
 32 (D) SOFTWARE: Winvatin (Genentech)

34 (vi) CURRENT APPLICATION DATA:

C--> 35 (A) APPLICATION NUMBER: US/09/717,641  
 C--> 36 (B) FILING DATE: 21-Nov-2000  
 37 (C) CLASSIFICATION:

67 (vii) PRIOR APPLICATION DATA:

40 (A) APPLICATION NUMBER: 08/922345  
 41 (B) FILING DATE: 03-SEP-1997  
 44 (A) APPLICATION NUMBER: 08/463587  
 45 (B) FILING DATE: 05-JUN-1995  
 48 (A) APPLICATION NUMBER: 08/050058  
 49 (B) FILING DATE: 30-APR-1993  
 52 (A) APPLICATION NUMBER: PCT/US91/09133  
 53 (B) FILING DATE: 03-DEC-1991  
 56 (A) APPLICATION NUMBER: 07/743614  
 57 (B) FILING DATE: 09-AUG-1991  
 60 (A) APPLICATION NUMBER: 07/715300  
 61 (B) FILING DATE: 14-JUN-1991  
 64 (A) APPLICATION NUMBER: 07/683400  
 65 (B) FILING DATE: 10-APR-1991  
 68 (A) APPLICATION NUMBER: 07/621667  
 69 (B) FILING DATE: 03-DEC-1990

71 (viii) ATTORNEY/AGENT INFORMATION:

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72      (A) NAME: Schwartz, Timothy R.
73      (B) REGISTRATION NUMBER: 32171
74      (C) REFERENCE/DOCKET NUMBER: P0645P4D2C3
76      (ix) TELECOMMUNICATION INFORMATION:
77      (A) TELEPHONE: 650/225-7467
78      (B) TELEFAX: 650/952-9881
79      (2) INFORMATION FOR SEQ ID NO: 1:
81      (i) SEQUENCE CHARACTERISTICS:
82      (A) LENGTH: 36 base pairs
83      (B) TYPE: Nucleic Acid
84      (C) STRANDEDNESS: Single
85      (D) TOPOLOGY: Linear
87      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
90      GCCAGCTGTG GCTTCTAGAG TGGCGGCGGC TCTGGT 36
92      (2) INFORMATION FOR SEQ ID NO: 2:
94      (i) SEQUENCE CHARACTERISTICS:
95      (A) LENGTH: 36 base pairs
96      (B) TYPE: Nucleic Acid
97      (C) STRANDEDNESS: Single
98      (D) TOPOLOGY: Linear
100     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
103     AGCTGIGGCT TCGGGCCCTT AGCATTTAAT GCGGTA 36
105     (2) INFORMATION FOR SEQ ID NO: 3:
107     (i) SEQUENCE CHARACTERISTICS:
108     (A) LENGTH: 33 base pairs
109     (B) TYPE: Nucleic Acid
110     (C) STRANDEDNESS: Single
111     (D) TOPOLOGY: Linear
113     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
116     TTCACAAACG AAGGGCCCTT AATTAAAGCC AGA 33
118     (2) INFORMATION FOR SEQ ID NO: 4:
120     (i) SEQUENCE CHARACTERISTICS:
121     (A) LENGTH: 30 base pairs
122     (B) TYPE: Nucleic Acid
123     (C) STRANDEDNESS: Single
124     (D) TOPOLOGY: Linear
126     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
129     CAATAATAAC GGGCTAGCCA AAAGAACTGG 30
131     (2) INFORMATION FOR SEQ ID NO: 5:
133     (i) SEQUENCE CHARACTERISTICS:
134     (A) LENGTH: 24 base pairs
135     (B) TYPE: Nucleic Acid
136     (C) STRANDEDNESS: Single
137     (D) TOPOLOGY: Linear
139     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
142     CACGACAGAA TTCCCGACTG GAAA 24
144     (2) INFORMATION FOR SEQ ID NO: 6:
146     (i) SEQUENCE CHARACTERISTICS:
147     (A) LENGTH: 23 base pairs

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148      (B) TYPE: Nucleic Acid
149      (C) STRANDEDNESS: Single
150      (D) TOPOLOGY: Linear
152      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
155      CTGTTTCTAG ACIGAAATTG TTA 23
157      (2) INFORMATION FOR SEQ ID NO: 7:
159      (i) SEQUENCE CHARACTERISTICS:
160          (A) LENGTH: 21 base pairs
161          (B) TYPE: Nucleic Acid
162          (C) STRANDEDNESS: Single
163          (D) TOPOLOGY: Linear
165      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
168      ACATTCCTGG GTACCGTCA G 21
170      (2) INFORMATION FOR SEQ ID NO: 8:
172      (i) SEQUENCE CHARACTERISTICS:
173          (A) LENGTH: 63 base pairs
174          (B) TYPE: Nucleic Acid
175          (C) STRANDEDNESS: Single
176          (D) TOPOLOGY: Linear
178      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
181      GCTTCAGGAA GGACATGGAC NNSGTCNNSA CANN SCTGNN SATCGTGCAG 50
183      TGCCGCTCTG TGG 63
185      (2) INFORMATION FOR SEQ ID NO: 9:
187      (i) SEQUENCE CHARACTERISTICS:
188          (A) LENGTH: 24 base pairs
189          (B) TYPE: Nucleic Acid
190          (C) STRANDEDNESS: Single
191          (D) TOPOLOGY: Linear
193      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
196      AAGGTCTCCA CATACCTGAG GATC 24
198      (2) INFORMATION FOR SEQ ID NO: 10:
200      (i) SEQUENCE CHARACTERISTICS:
201          (A) LENGTH: 33 base pairs
202          (B) TYPE: Nucleic Acid
203          (C) STRANDEDNESS: Single
204          (D) TOPOLOGY: Linear
206      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
209      ATGGACAAGG TCTCGACAFA CCTGCGCATC GTG 33
211      (2) INFORMATION FOR SEQ ID NO: 11:
213      (i) SEQUENCE CHARACTERISTICS:
214          (A) LENGTH: 36 base pairs
215          (B) TYPE: Nucleic Acid
216          (C) STRANDEDNESS: Single
217          (D) TOPOLOGY: Linear
219      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
222      GGCAGCTGTG GATTCTACAG TGGCGGTGGC TCTGGT 36
224      (2) INFORMATION FOR SEQ ID NO: 12:
226      (i) SEQUENCE CHARACTERISTICS:
227          (A) LENGTH: 12 amino acids

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228      (B) TYPE: Amino Acid
229      (D) TOPOLOGY: Linear
231      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
233      Gly Ser Cys Gly Phe Glu Ser Gly Gly Ser Gly
234      1          5          10
236 (2) INFORMATION FOR SEQ ID NO: 13:
238      (i) SEQUENCE CHARACTERISTICS:
239          (A) LENGTH: 27 base pairs
240          (B) TYPE: Nucleic Acid
241          (C) STRANDEDNESS: Single
242          (D) TOPOLOGY: Linear
244      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
247      CGGACTGGGC AGATAFICAA GCAGACC 27
249 (2) INFORMATION FOR SEQ ID NO: 14:
251      (i) SEQUENCE CHARACTERISTICS:
252          (A) LENGTH: 38 base pairs
253          (B) TYPE: Nucleic Acid
254          (C) STRANDEDNESS: Single
255          (D) TOPOLOGY: Linear
257      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:
260      CTCGAAGAACT ACGGGTTACC CTGACTGCTT CAGGAAGG 38
262 (2) INFORMATION FOR SEQ ID NO: 15:
264      (i) SEQUENCE CHARACTERISTICS:
265          (A) LENGTH: 30 base pairs
266          (B) TYPE: Nucleic Acid
267          (C) STRANDEDNESS: Single
268          (D) TOPOLOGY: Linear
270      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:
273      CGCATCGIGC AGTGCAGAIC TGTGGAGGGC 30
275 (2) INFORMATION FOR SEQ ID NO: 16:
277      (i) SEQUENCE CHARACTERISTICS:
278          (A) LENGTH: 66 base pairs
279          (B) TYPE: Nucleic Acid
280          (C) STRANDEDNESS: Single
281          (D) TOPOLOGY: Linear
283      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:
286      GTTACTCTAC TGCTTTCAGG AAGGACATGG ACNNSGTCNN SACANNSCTG 50
288      NNSATCGTGC ACTGCA 66
290 (2) INFORMATION FOR SEQ ID NO: 17:
292      (i) SEQUENCE CHARACTERISTICS:
293          (A) LENGTH: 64 base pairs
294          (B) TYPE: Nucleic Acid
295          (C) STRANDEDNESS: Single
296          (D) TOPOLOGY: Linear
298      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:
301      GATCTGCACT GCACGATSNN CAGSNNTGTS NNGACSNNGT CCATGTCCTT 50
303      CCIGAAGCAC TAGA 64
305 (2) INFORMATION FOR SEQ ID NO: 18:
307      (i) SEQUENCE CHARACTERISTICS:

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308      (A) LENGTH: 25 base pairs
309      (B) TYPE: Nucleic Acid
310      (C) STRANDEDNESS: Single
311      (D) TOPOLOGY: Linear
313      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
316      GCCITTGACA GGTACCAGGA GTTIG 25
318      (2) INFORMATION FOR SEQ ID NO: 19:
320      (i) SEQUENCE CHARACTERISTICS:
321      (A) LENGTH: 33 base pairs
322      (B) TYPE: Nucleic Acid
323      (C) STRANDEDNESS: Single
324      (D) TOPOLOGY: Linear
326      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
329      CCAACIATAC CACTCTCGAG GTCTATTGTA TAA 33
331      (2) INFORMATION FOR SEQ ID NO: 20:
333      (i) SEQUENCE CHARACTERISTICS:
334      (A) LENGTH: 66 base pairs
335      (B) TYPE: Nucleic Acid
336      (C) STRANDEDNESS: Single
337      (D) TOPOLOGY: Linear
339      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
342      TCGAGGCTCN NSGACAACGC GNNSCIGCGT GCTNNSCGTC TTNNSCAGCT 50
344      GGCCITTGAC ACGTAC 66
346      (2) INFORMATION FOR SEQ ID NO: 21:
348      (i) SEQUENCE CHARACTERISTICS:
349      (A) LENGTH: 58 base pairs
350      (B) TYPE: Nucleic Acid
351      (C) STRANDEDNESS: Single
352      (D) TOPOLOGY: Linear
354      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:
357      GTGTCAAAGG CCAGCTGSNN AAGACGSNNA GCACGCAGSN NCGCGTTGTC 50
359      SNNAGGCC 58
361      (2) INFORMATION FOR SEQ ID NO: 22:
363      (i) SEQUENCE CHARACTERISTICS:
364      (A) LENGTH: 65 base pairs
365      (B) TYPE: Nucleic Acid
366      (C) STRANDEDNESS: Single
367      (D) TOPOLOGY: Linear
369      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:
372      GTTACTCTAC TGCTTCNNSA AGGACATGNN SAAGGTCAGC NNSTACCTGC 50
374      GCNNSGTGCA GTGCA 65
376      (2) INFORMATION FOR SEQ ID NO: 23:
378      (i) SEQUENCE CHARACTERISTICS:
379      (A) LENGTH: 64 base pairs
380      (B) TYPE: Nucleic Acid
381      (C) STRANDEDNESS: Single
382      (D) TOPOLOGY: Linear
384      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:
387      GATCTGCACT GCACSNNGCG CAGGTASNNG CTGACCTTSN NCATGTCCTT 50

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VERIFICATION SUMMARY

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Input Set : A:\P0645P4D2C3.txt

Output Set: N:\CRF3\12062000\I717641.raw

L:3 M:244 W: Invalid beginning of sequence listing, Data=[SEQUENCE LISTING], Duplicate Sequence Listing Title:  
L:35 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]  
L:36 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]